SEQUENCE LISTING

General Information

APPLICANT: CSIR

TITLE OF INVESTIGATION: A Promoter For High-Throughput Screening For Inhibitors Against Mycobacteria Under Low Carbon Conditions

NUMBER OF SEQUENCES : 2

CORRESPONDENCE ADDRESS : Indian Institute of Science, Banglore, India

INFORMATION FOR SEQ ID No:1

1 SEQUENCE CHARACTERISTICS :

LENGTH : 1.5 kb.

TYPE : DNA

GGATCCCTGGTAACCCCCGACAGCTACCAGCGCACCGACTACCCGTCGGCCGGGATCGAGCAGCTGATCT
TCGCACCACAAGGTTCACTCGCGCAAAGCCGCACCCGCCGCGCGCTCGCGTTGTGTGTACCCCGGGACGC
GATCGCTCGGGATGCCGGGGTTCCGATTGCCAACTCGCGGCTGTCCCCGGCGACCGACGATGCCCTCACC
GATGCCGACGGCGCCGCCGAAGCACGTCAGTTCGGCCGGGTGGACCCCGCCGCCGCTCGCGACGCGCTG
GGTGGTACGCCGCTGACCGTGCGGATCGGCTACGGCAGGCCCAACGCTCGGTTGGCGGCCACCATCGGA
ACCATTGCCGACGCCTGCGCCCCGGCCGGGATCACCGTTTCGGATGTGACGGTGGACACACCCGGACCGC
AAGCGCTGCGGGACGGAAAGATTGACGTATTGTTGGCGAGCACCGGTGGGGCCACCGGCAGCGGATCGA
GCGGATCGTGTGCGATGGATGCCTATGACTTGCACAGCGGCAACGGAAACAATCTATCGGGGTACGCAA
ACGCTCAGATCGACGGCATCATCAGCGCGCTCGCGGTGTCGGCCGACCCCGCCGAGCGGGCCAGGTTGC
TTGCCGAGGCCGCGCCGGTGCTCTGGGATGAGATGCCAACCTTGCCGTTGTACCGGCAGCAGCGCACGTT
GTTGATGTCGACGAAAATGTATGCGGTGAGCAGGAATCCGACGCGATGGGGGGCAGGGTGGAACATGGA
TCGCTGGGCGCTGGCGCGGTGACGATGGCCAGTGCCATCTGCAGGTAATTGACAGAATTCCACGACGAG
AAGCGGACTATCGGAGCGTAGTGTCGCAGGTGCTCCGGGCTGTCTGGGAGAGGATGTGTGCCATGGCGG

TACATGGGCTGGTGACTACGTGTTGAACGTGATCGCGACGGGGCTCTCCTTAAAGGCACGGGGGAAGCG
CCGCCGGCAGCGTTGGGTCGACGACGGGCGGGTATTGGCGCTCGGTGAGTCCCGCCGGAGCTCAGCCAT
ATCTGTGGCCGACGTGGTTGCGTCGCTGACCCGGGATGTGGCCGACTTTCCGGTTCCCGGCGTCGAGTTC
AAGGACCTCACCCCGCTATTCGCCGACCGAAGAGGATTGGCCGCGGTAACCGAAGCGCTGGCCGATCGG
GCGTCCGGAGCTGACCTGGTGGCCGGCGTCGACGCCCGCGGGTTTCTGGTGGCAGCCGCGGTCGCCACCC
GGCTCGAAGTGGGTGTGCTGGCCGTTCGCAAGGGCGGCAAGCTGCCCCGGCCGGTGCTCAGCGAGGAGT
ACTACAGGGCGTACGGCGCCGCCACTCTGGAGATTCTCGCTGAGGGCATCGAGGTTGCGGGCCGCCGTGT
CGTGATCATTGACGACGTGTTAGCAACCGGCGGCACCATCGGCGCGACGCGACGCCTGCTTGAGCGCGG
TGGCGCCAACGTGGCCGGGGCGGCCGTAGTGGTGGAACTTGCGGGGTTGAGCGGTCGCGCGGCGCTCGC
ACCGCTGCCGGTGCACAGCCTGAGCCGCCTGTGAGGGATATCCTCTAGGTCGGAGGTGACGAACGTGGC
CGAGGACCAGCTCACGGCGCAAGCGGTTGCACCGCCCACGGAGGCTTCTGCGGCTCTCGAGCCCGCTCTC
GAGACGCCCGAGTCGCCGGTCGAGACTCTTAAGACCAGCATCAGCGCGTCGCGTCGGGTGCGGGCCCGA
TTGGCCCGGCGGATGACCGCCCAGCGCAGCACCACCAATCCGGTGCTCGAGCCGTTGGTGGCGGTGCAC
CGGGAGATCTATCCCAAGGCCGACCTGTCGATCTTGCAGCGAGCCTACGAGGTCGCTGACCAAAGGCAT
GC

ORGANISM : *M.tuberculosis*

IMMEDIATE : Natural

NAME / KEY : Natural oligonucleotide

SEQUENCE ID # 1

INFORMATION FOR SEQ ID No:2

1 SEQUENCE CHARACTERISTICS :

LENGTH : 200 bp

TYPE : DNA

Sak2 ⟶

CGCCGCCACTCTGGAGATTCTCGCTGAGGGCATCGAGGTTGCGGGCCGCCGTGTCGTGATCATTGACGAC

GTGTTAGCAACCGGCGGCACCATCGGCGCGACGCGACGCCTGCTTGAGCGCGGTGGCGCCAACGTGGCC

GGGGCGGCCGTAGTGGTGGAACTTGCGGGGTTGAGCGGTCGCGCGGCGCTCGCACCGCTGCCGGTGCAC

⟵ Sak1

AGCCTGAGCCGCCTGTGAGGGA**TATCCT**CTAGGTCGGAGGTGACGAAC**GTG**GCCGAGG

ORGANISM : Natural sequence

IMMEDIATE : Natural

NAME / KEY : Natural oligonucleotide

SEQUENCE ID # 2

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